1/32

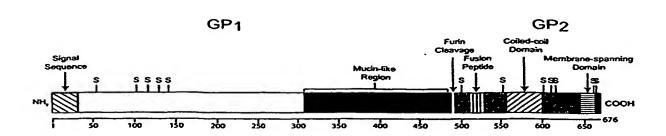


fig. 1A

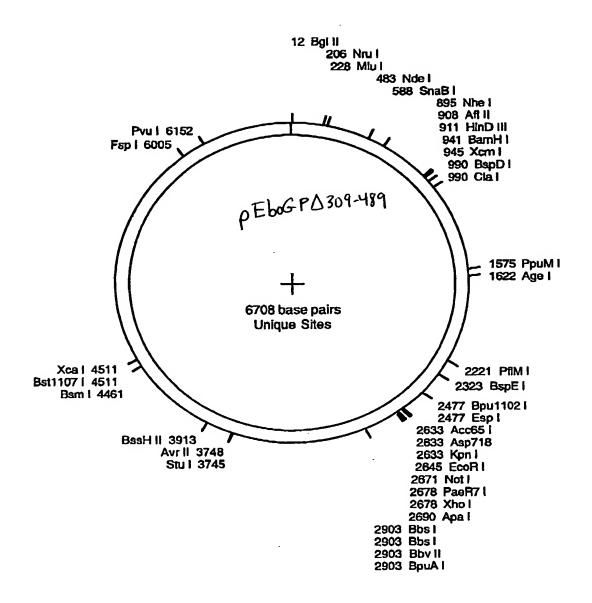


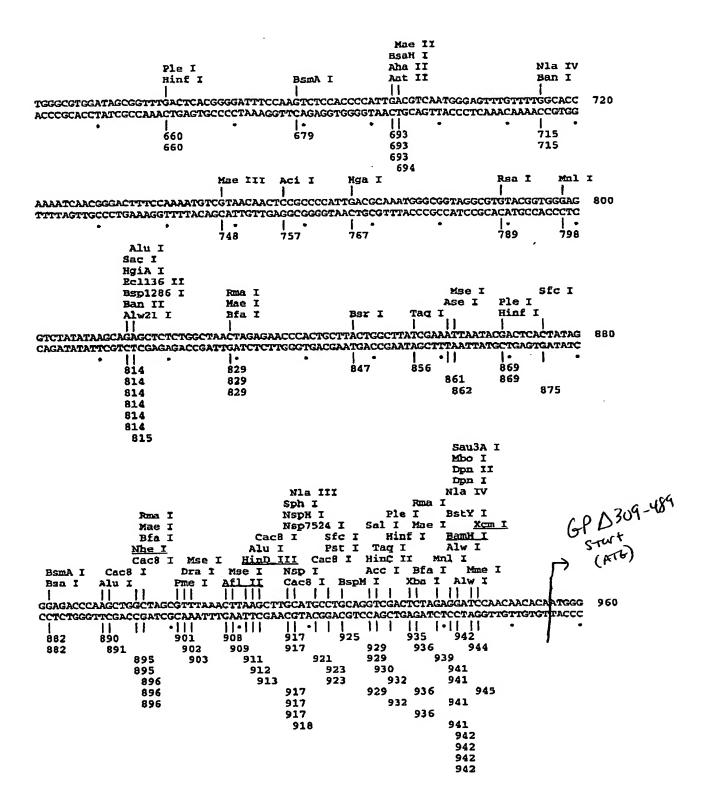
fig. 1B

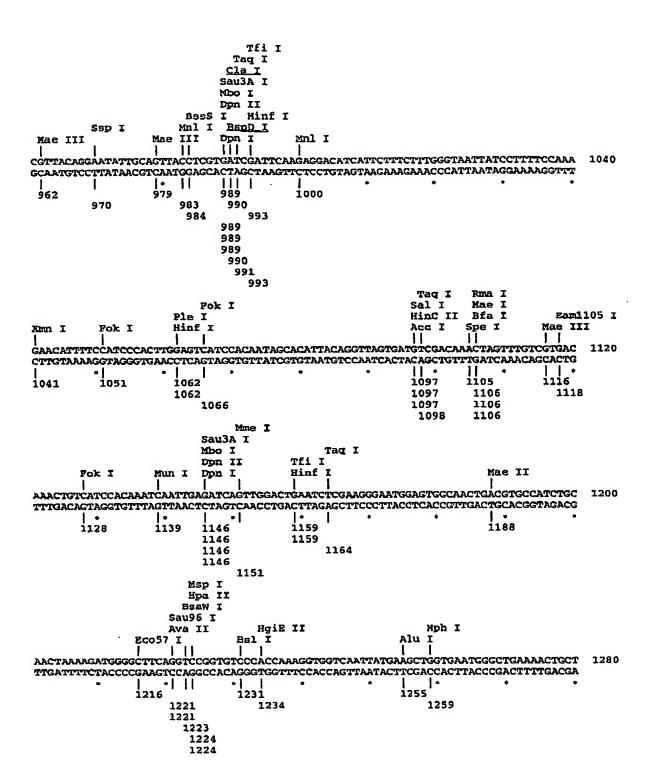
Positions of Restriction Endonucleases sites (unique sites underlined)

3/32

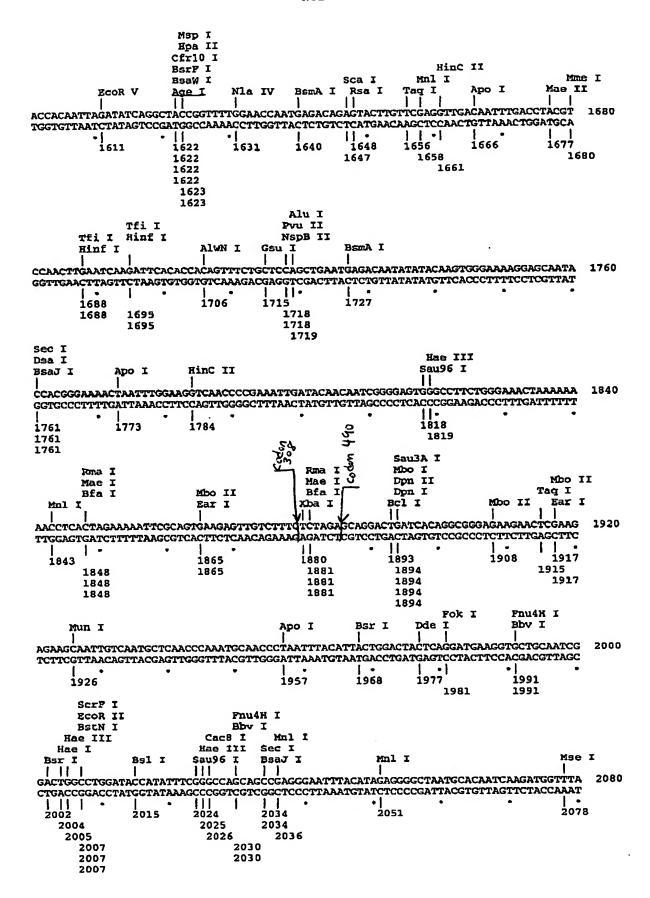
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                   Sau3A I
                                 Hinf I
   Mbo I
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   Dpn II Dpn I
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                                        Rsa I
                                                         Fnu4H I
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GACGGATCGGGÁGATCTCCCGATCCCCTATGGTCGACTCTCAGTACAATCTGCTCTGATGCCGCATAGTTAAGCCAGTAT
CTGCCTAGCCCTCTAGAGGGCTAGGGGATACCAGCTGAGAGTCATGTTAGACGAGACTACGGCGTATCAATTCGGTCATA
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       Tth111 II
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CTGCTCCTGCTTGTGTGTTGGAGGTCGCTGAGTAGTGCGCGAGCAAAATTTAAGCTACAACAAGGCAAGGCTTGACCGA 160
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                                                       Hae III
                                                                  Bstu I
     Nla III
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                                             BstU I
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                                                                 Mlu I
Mun I
        Mbo II
                  Dde I
                                    Hha I
                                            Nru I Rsa I
                                                                 Afl III
CAATTGCATGAAGAATCTGCTTAGGGTTAGGCGTTTTGCGCTGCTTCGCGATGTACGGGGCCAGATATÁCGCGTTGACATT
GTTAACGTACTTCTTAGACGAATCCCAATCCGCAAAACGCGACGAAGCGCTACATGCCCGGTCTATATGCGCAACTGTAA
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           173
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        Rma I
        Mae I
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        Bfa I Mse I
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        Spe I Ase I
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GATTATTGÁCTAGTTÁÍTAATAGTAATCAATTACGGGGTCATTAGTTCATAGCCCATATATGGAGTTCCGCCTTACATAA 320
CTAATAACTGATCAATAATTATCATTAGTTAATGCCCCCAGTAATCAAGTATCGGGTATATACCTCAAGGCGCAATGTATT
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                      Aha II
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                                                                 Rsa I
                      Aat II
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TTGCGGTTATCCCTGAAAGGTAACTGCAGTTACCCACCTGATAAATGCCATTTGACGGGTGAACCGTCATGTAGTTCACA
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                                             Bgl I
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Aat II
                                                                 Rsa I
                                            Sau96 I
                                            Hae III
                                                               Bsr I
  Nde_I
            Rsa I
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 ACCCTGAAAGGATGAACCGTCATGTAGATGCATAATCAGTAGCGATAATGGTACCACTACGCCAAAACCGTCATGTAGTT
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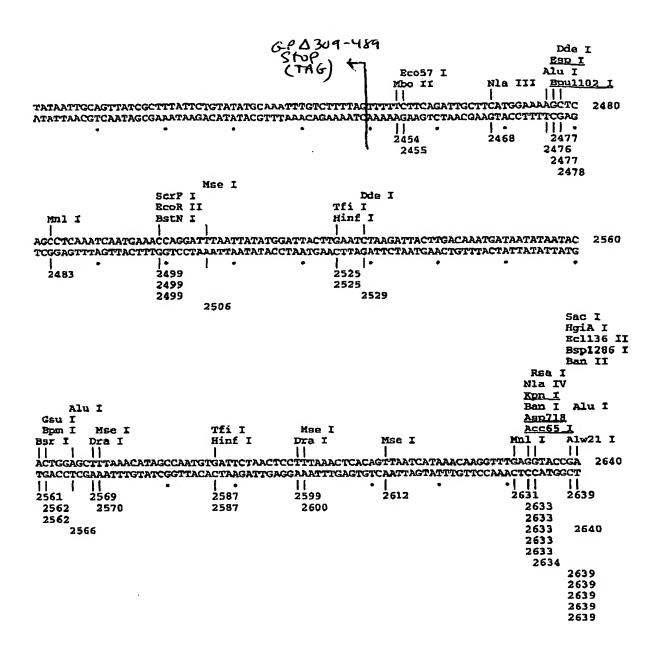




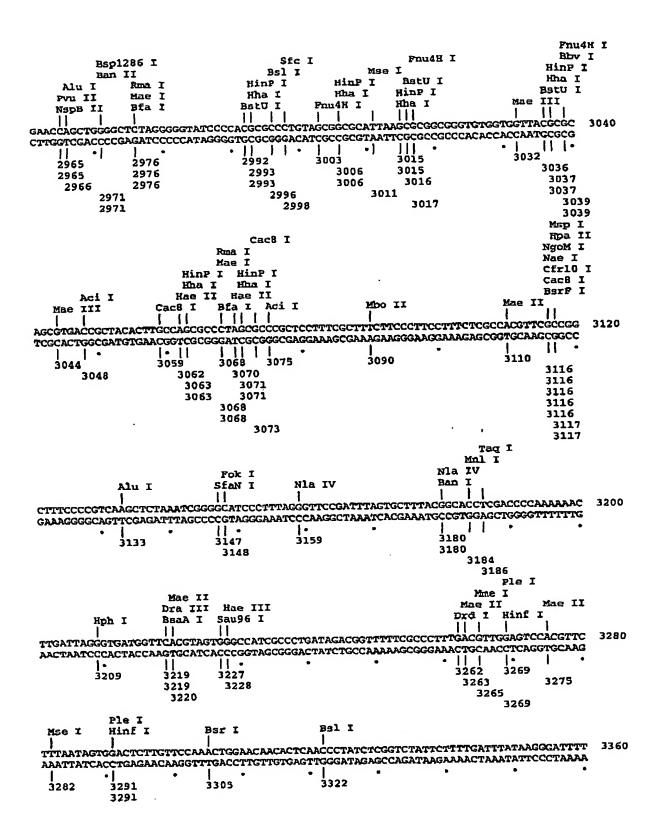
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ACAATCTTGAAATCAAAAAACCTGACGGGAGTGAGTGTCTACCAGCAGCGCCAGACGGGATTCGGGGGTTCCCCCGGTGC
TGTTAGAACTTTAGTTTTTTGGACTGCCCTCACTCACAGATGGTCGTCGCGGTCTGCCCTAAGCCCCGAAGGGGGCCCACG
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      HgiA I
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                                       Bama I
      Bsp1286 I
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                                     Map I
      ApaL I
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                                                     Bal I Mnl I
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      Alw21 I
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CCCATACACGTGTTTCATAGTCCTTGCCCTGGCACACGCCTCTGAAACGGAAGGTATTTCTCCCACGAAAGAAGGACAT
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 Sau3A I
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 Don II
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                           BsaJ I
                                            Eco57 I
 Dpn I
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                                      BsrF I
             Mbo II
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 GATTCTTCCTGAAGAAGTCGAGTGTGGGGAACTCTCTCGGCCAGTTACGTTGCCTCCTGGGCAGATCACCGATGATAAGA
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GAAGGCCTGGTCCCCCTGTTACTGTTAACCACCTGTCCTACCTCTGTTACCTATGGCCGTCCATAACCTCAATGTCCGCA
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GCTCGAATTCTGCAGATATCCAGCACAGTGGCGGCCGCTCGAGTCTAGAGGGCCCGTTTAAACCCGCTGATCAGCCTCGA 2720
CGAGCTTAAGACGTCTATAGGTCGTGTCACCGCCGGCGAGCTCAGATCTCCCGGGCAAATTTGGGCGACTAGTCGGAGCT
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GTCCTTTCCTAATAAAATGAGGAAATTGCATCGCATTGTCTGAGTAGGTGTCATTCTATTCTGGGGGGTGGGGTGGGGCA 2880
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CCTGTCGTTCCCCCTCCTAACCCTTCTGTTATCGTCCGTACGACCCCTACGCCACCCGAGATACCGAAGACTCCGCCTTT
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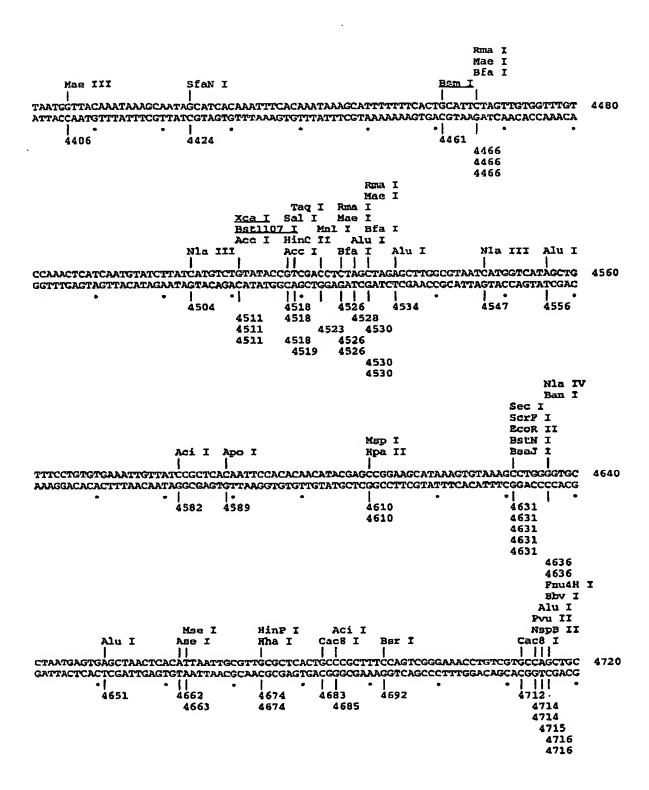
WO 03/102219

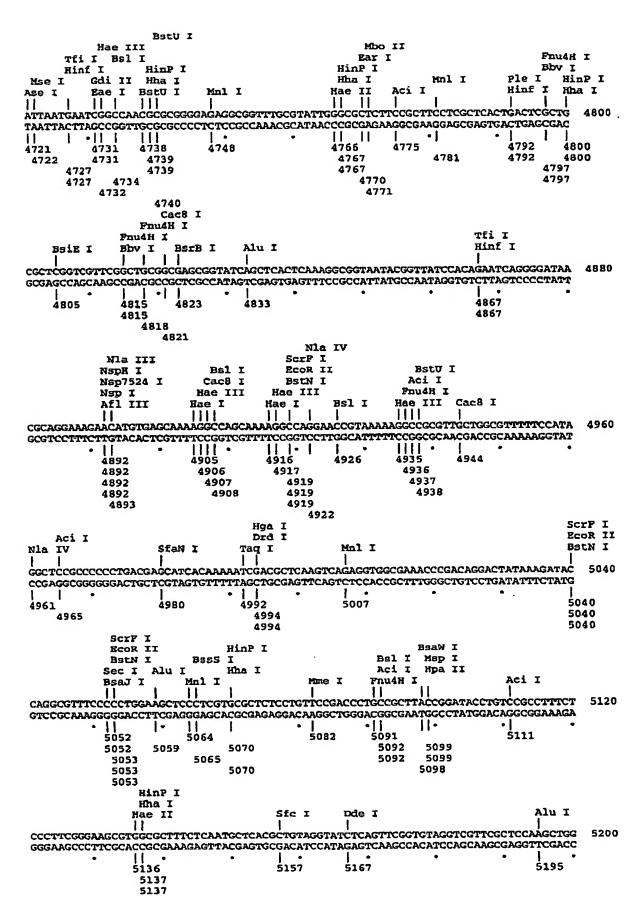
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BsiE I BsrB I Msp I Hpa II Taq I GTCGCCGGAGCGGTCGAGTCAGCGGCCTCGCCAGCTCAGC	AGYCCLCCCLCCCCGYCCC	ScrP I Nci I Msp I Hpa II Bcn I Sma I Sec I ScrF I Nci I BaaJ I Bcn I Ava I GTTCCCCGGGACTTCC CAAGAGGGCCCTGAAGG 1965 3965 3965 3965 3965 3965 3966 3966 3	acctcctgctgaa	Msp I Sau96 I Rpa II Cfr10 I BSIF I Ava II GCCCGGTGTGG 4000 GCGGCCACACC • • 3991 3999 3991 3992 3992 3999

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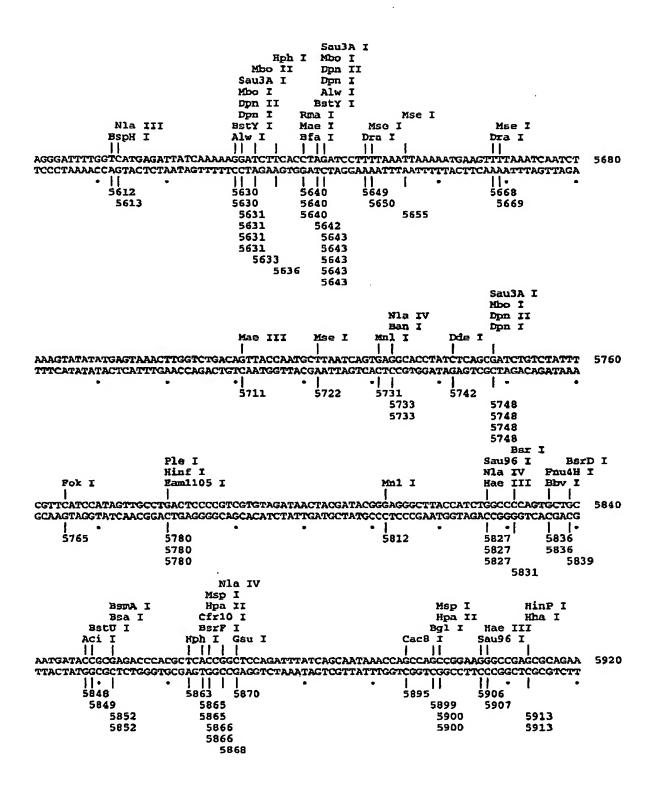
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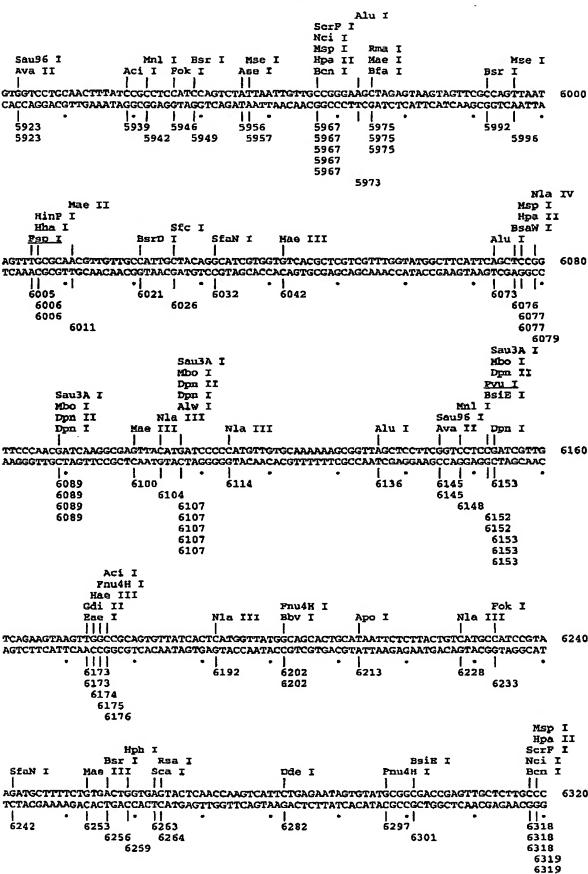




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HinP I
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    HgiA I
                              Bbv I
                                            Mae III
    Bsp1286 I
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    Apal I
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    Alw21 I
                          BsiE I Hha I
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CGACACACGTGCTTGGGGGCCAAGTCGGGCTGGCGACGCGGAATAGGCCATTGATAGCAGAACTCAGGTTGGGCCATTCT
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CACGACTTATCGCCÁCTGGCAGCAGCCÁCTGGTAACAGGATTAGCAGAGCGAGGTATGTAGGCGGTGCTACAGAGTTCTT 5360
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GAAGTGGTGGCCTAACTACGGCTACACTAGAAGGACAGTATTTGGTATCTGCGCTCTGCTGAAGCCAGTTACCTTCGGAA 5440
CTTCACCACCGGATTGATGCCGATGTGATCTTCCTGTCATAAACCATAGACGCGAGACGACTTCGGTCAATGGAAGCCTT
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<u>ANAGAGTTGGTÁGCTCTTGATCCGGCANACANÀCCACCGCTGGTAGCGGTGGTTTTTTTGTTTĞCANGCAGCAGATTACĞ</u>
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CGCAGAAAAAAGGATCTCAAGAAGATCCTTTGATCTTTTCTACGGGGTCTGACGCTCAGTGGAACGAAAACTCACGTTA
GCGTCTTTTTTCCTAGAGTTCTTCTAGGAAACTAGAAAAGATGCCCCAGACTGCGAGTCACCTTGCTTTTGAGTGCAAT
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22/32



ACC I

Acc65 I

Bsp1286 I

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Psp I

PCT/US03/17577

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Hinp I
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                                               Bsp1286 I
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GGCGTCAATACGGGATAATACCGCGCCACATAGCAGAACTTTAAAAGTGCTCATCATTGGAAAACGTTCTTCGGGGCGAA 6400
CCGCAGTTATGCCCTATTATGCCGCGGTGTATCGTCTTGAAATTTTCACGAGTAGTAACCTTTTGCAAGAAGCCCCGCTT
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AACTCTCAAGGATCTTACCGCTGTTGAGATCCAGTTCGATGTAACCCACTCGTGCACCCAACTGATCTTCAGCATCTTTT
TTGAGAGTTCCTAGAATGGCGACAACTCTAGGTCAAGCTACATTGGGTGAGCACGTGGGTTGACTAGAAGTCGTAGAAAA
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ACTITICACCAGCGTTTCTGGGTGAGCAAAAACAGGAAGGCAAAATGCCGCAAAAAAGGGAATAAGGGCAACACGGAAATG
TGARAGTGGTCGCARAGACCCACTCGTTTTTGTCCTTCCGTTTTACGGCGTTTTTTCCCTTATTCCCGCTGTGCCTTTAC
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TTGAATACTCATACTCTTTTTCAATATTATTGAAGCATTTATCAGGGTTATTCTCACAGCGGATACATATTTG 6640
AACTTATGAGTATGAGAAGGAAAAGTTATAATAACTTCGTAAATAGTCCCAATAACAGAGTACTCGCCTATGTATAAAC
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 AATGTATTTAGAAAAATAAACAAATAGGGGTTCCGCGCACATTTCCCCGAAAAGTGCCACCTGACGTC
 TTACATAAATCTTTTTATTTGTTTATCCCCAAGGCGCGTGTAAAGGGGCTTTTCACGGTGGACTGCAG
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      Restriction Endonucleases site usage
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NspH I

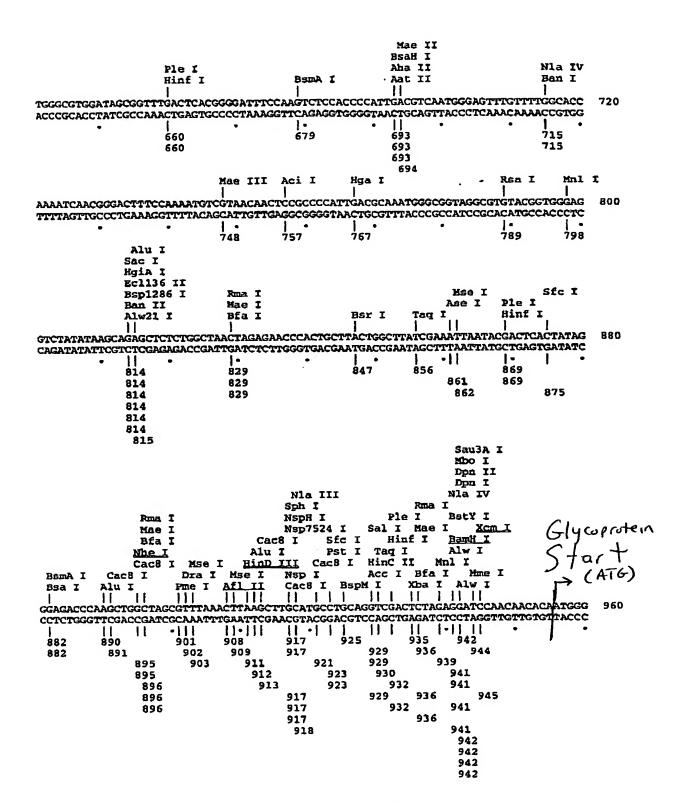


fig. 1C

WO 03/102219

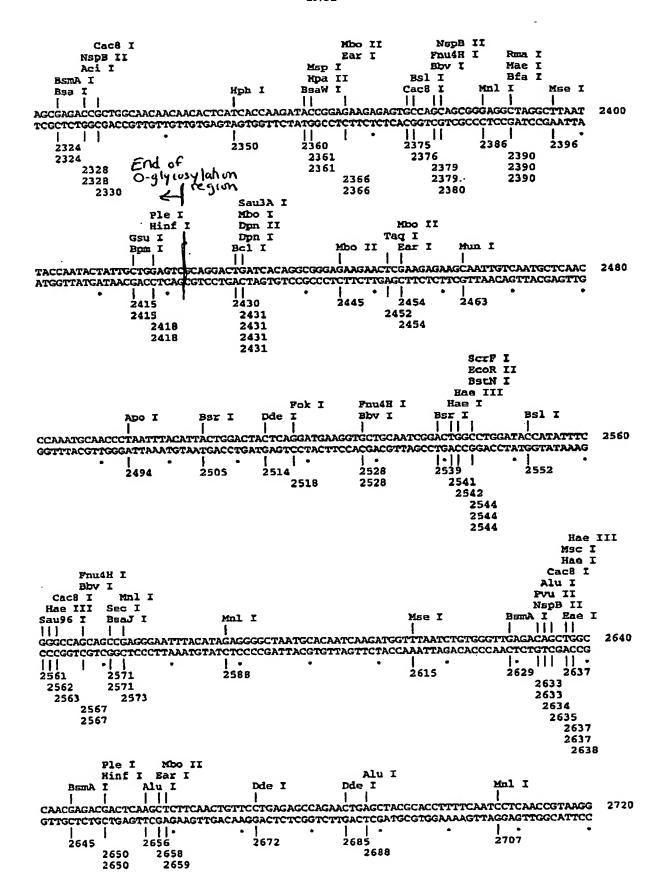
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GCCATACACGTGTTTCATAGTCCTTGCCCTGGCACACGGCCTCTGAAACGGAAGCTATTTCTCCCACGAAAGAAGAAGAACAT
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 Mbo I
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 Dpn II
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                                              Eco57 I
                                                                                  Alu I
 Don I
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ACTAGCTGAACGA,AGGTGTCAATAGATGCCTCCTTGCTGAAAGCGACTTCCACAGCAACGTAAAGACTATGACGGGGTTC
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CATTCTTCCTGAAGAAGTCGAGTGTGGGGAACTCTCTCGGCCAGTTACGTTGCCTCCTGGGCAGATCACCGATGATAAGA
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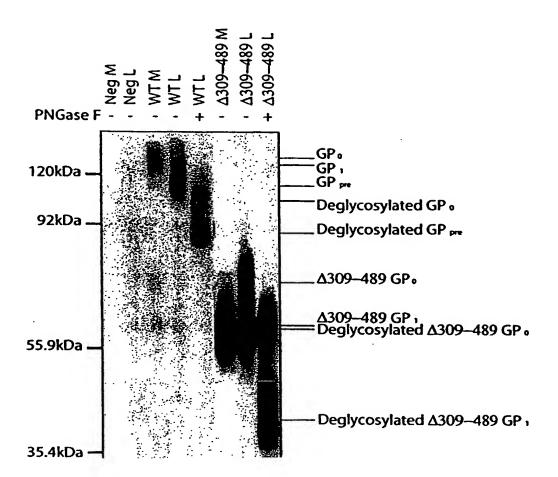
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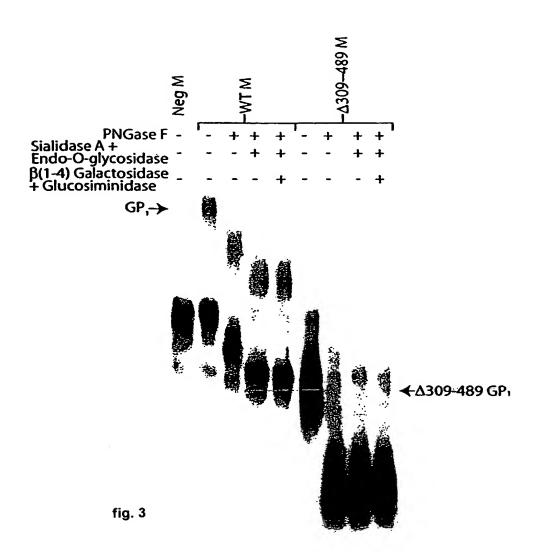


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TGGTTCTTGTATTGTCTGTTTTAACTAGTCTAATAAGTACTAAAACAACTATTTTGGGAAGGCCTGGTCCCCCTGTTACT
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fig. 2



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